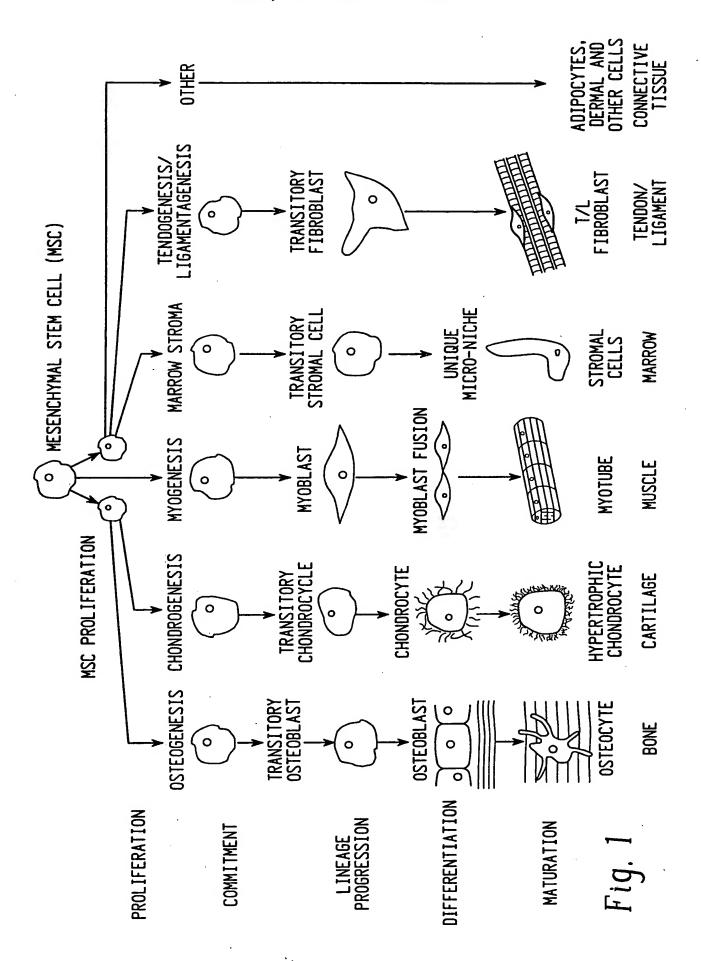
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Fig. 2

CHONDROGENIC LINEAGE

COMMITTED Chondroprogenitor : Type I Collagen

Chondrogenic PROGENITOR : Type VI Collagen, CSPG-M

Chondroblast : Type II Collagen, CDI(28Kd),

CSPG-H (aggrecan)

Chondrocyte I : 148Kd, link, protein

↓Hypertrophic Chondrocyte : Type X Collagen, 59Kd,

Chondrocyte II

 $1,25-(OH)_2D_3R$

Calcifying Chondrocyte : Type I Collagen, 63Kd, osteopontin,

osteocalin, steonectin

100 Kd, matric, vesicle production

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Fig. 3A

Sequence of CZF-1 (cDNA)
50 AATGGAGCGAAGACCATGGGGACTGAGTACACAGATGAAGACACAGAAGC
100 ATAGAGAGGATAAGTAATCACTAGCAAGTGGAAGAACCGGGATTCAGATC
150 CAGAACAGGCTGACTCCAGAGTCACTGGCTGTCATGTAGTTTCCTCAACT
200 ACTGCCTCAGCTCTACAATCCCAGAGTAAAGCTCTTCTCCAAATGAAGAG
250 CCAGGAAGAGGTAGAGGTGGCAGGAATTAAACTTTGTAAAGCCATGTCCC
300 TGGGTTCACTGACTTTCACAGATGTGGCCATAGACTTTTCCCAAGATGAA
350 TGGGAGTGGCTGAATCTTGCTCAGAGAAGTTTGTACAAGAAGGTGATGTT
400 AGAAAACTACAGGAACCTAGTTTCAGTGGGTCTTTGCATTTCTAAACCAG
450 ATGTGATCTCCTTACTGGAGCAAGAGAAAGACCCTTGGGTGATAAAAGGA
500 GGGATGAACAGAGGCCTGTGCCCAGACTTGGAGTGTGTGT
550 ATCATTATCTTTAAACCAGGATATTTATGAAGAAAAATTACCCCCGGCAA
600 TCATAATGGAAAGACTTAAAAGCTATGACCTTGAATGTTCAACATTAGGG
650 AAAAACTGGAAATGTGAAGACTTGTTTGAGAGGGAGCTTGTAAACCAGAA
700 GACACATTTTAGGCAAGAGACCATCACTCATATAGATACTCTTATTGAAA
750 AAAGAGATCACTCTAACAAATCTGGGACAGTTTTTCATCTGAATACATTA
. 800

TCTTATATAAAACAGATTTTTCCCATGGAAGAGAGAATATTTAATTTTCA

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Fig. 3B

850 TACAGATAAGAAAAGCTTAAAAACACATTCAGTTGTGAAAAAAACACAAGC
900 AAGACCGTGGAGAAAAGAAACTTTTAAAATGTAATGACTGTGAGAAAATA
950
TTCAGCAAAATCTCAACCCTTACTCTTCACCAAAGAATTCATACAGGAGA
1000
GAAACCCTATGAATGTATTGAATGTGGAAAGGCCTTTAGCCAGAGTGCCC
1050
ACCTTGCTCAACATCAGAGAATACACACAGGAGAAAAACCTTTTGAATGT
1100
ACTGAATGTGGGAAAGCCTTCAGCCAGAATGCTCATCTTGTTCAACACCA
1150
GAGAGTTCATACTGGAGAGAAACCTTATCAGTGTAAGCAGTGTAATAAAG
1200
CATTCAGCCAGCTTGCACACCTTGCTCAACATCAGAGGGTCCACACTGGA
1250
GAGAAACCCTATGAATGTATTGAATGTGGGAAGGCTTTTAGTGATTGCTC
1300
ATCCCTAGCTCATCGAAGGATTCACACTGGGAAAAGACCTTATGAAT
1350
GTATTGACTGTGGGAAAGCTTTCAGGCAGAATGCTTCTCTTATACGTCAT
1400 CGGCGATATTATCATACTGGAGAGAAACCCTTTGACTGTATTGATTG
CGGCGATATTATCATACTGGAGAGAAACCCTTTGACTGTATTGATTG
1450 GAAGGCTTTCACTGATCACATAGGACTTATTCAGCATAAGAGAATTCATA
GAAGGC111CAC1GA1CACA1AGGAC11A11CAGCA1AAGAGAA11CA1A
1500 CTGGAGAGAGACCTTACAAATGTAATGTGTGTGGGAAGGCTTTTAGCCAT
CTGGAGAGACCTTACAAATGTAATGTGTGTGGGAAGGCTTTTAGCCAT
1550 GGCTCATCTCTGACAGTACATCAGAGAATTCATACAGGAGAGAAACCTTA
GGC CA C C GACAG ACA CAGAGAA CA ACAGGAGAGAAACC IA
1600 TGAATGCAATATCTGTGAGAAAGCCTTCAGCCATCGTGGGTCTCTTACTC
IGMAIGCMAIMICIGIGAAMAGCCIICAGCCAICGIGGGICICIIACIC

Fig. 3C

1650
TTCATCAGAGAGTTCATACTGGAGAGAAACCCTATGAATGTAAAGAATGT
1700
GGGAAAGCTTTCCGGCAGAGCACGCATCTGGCTCATCATCAGAGAATTCA
1750
TACTGGAGAGAAACCTTATGAATGTAAGGAATGCAGCAAAACCTTCAGCC
1800
AGAATGCACACCTCGCGCAGCATCAGAAAATACACACTGGGGAGAAGCCT
1850
TATGAATGTAAGGAACGTGGTAAGGCCTTCAGTCAGATTGCACACCTTGT
1900
TCAGCACCAGAGAGTTCATACTGGTGAGAAGCCTTACGAATGTATTGAAT
1950
GTGGGAAGGCCTTTAGTGATGGCTCATATCTTGTTCAACATCCGAGACTC
2000
CACAGTGGCAAAAGACCGTATGAATGTCTTGAATGTGGGAAGGCATTCAG
2050 GCAGAGGGCATCCTTGATTTGTCATCAGAGATGTCATACTGGTGAGAAAC
GCAGAGGCATCCTTGATTTGTCATCAGAGATGTCATACTGGTGAGAAAC
2100
CTTATGAATGTAATGTTTGTGGGAAAGCCTTTAGCCATCGTAAATCCCTT
2150 ACTCTGCATCAGAGAATTCATACAGGAGAGAAACCTTATGAGTGTAAGGA
ACTCTGCATCAGAGAATTCATACAGGAGAGAAACCTTATGAGTGTAAGGA
2200 ATGTAGCAAAGCCTTCAGCCAGGTTGCCCATCTTACTCTACATAAGAGAA
ATOTAGCAAAGCCTTCAGCCAGGTTGCCCATCTTACTCTACATAAGAGAA
2250 TTCATACTGGAGAAAGGCCCTATGAGTGTAAAGAATGTGGAAAAGCCTTC
TICHIMCIOCACAAGGCCCIAIGAGIGIAAAGAAIGIGGAAAAGCCIIC
2300 AGGCAGAGTGTACATCTTGCTCATCATCAGCGAATTCATACCGGAGAGTC
2350 ATCAGTTATTCTCTCTCTCCCCCCCATACCACCAAGTCCTATAGATTC
2400 AATCTCGTAAATGCTTCTAGCATCCATCTGCTTTTTTCCAGCACATGTCC

Fig. 3D

2450 CATCATCATAGTCCAAGACGCAACCATCTCATCTGGATTTCTGCAGTAGC
2500 ATAACTGTTGCCCCTTTTGCTTCTATCAACTACATGTTTAACACTGTAGG
2550 CAGCCTAACCTTTTAAAAATAAAATACATAATTTATGTTATTTTCCCAT
2600 TTAAAACACTTGATTTGAAAAATATATTAACTAATCCATTTCAAGGATTT
2650 AGCACACACTGGCATATAGTTATTGCTAAATAAATGCTAGCCATTAAGGT
2666 AAAAAAAAAAAAA

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KRAB-A KRAB-B MKSQEEVEVAGIKLCKAMSLGSLTFTDVAIDFSQDEWEWLNLAQRSLYKKVMLENYRNLVSVGLCISKPDVISLLEQ EKDPW/IKGGMNRGLCPDLECVW/TKSLSLNQDIYEEKLPPAIIMERLKSYDLECSTLGKNWKCEDLFERELVNQKT HFRQETITHIDTLIEKRDHSNKSGTVFHLNTLSYIKQIFPMEERIFNFHTDKKSLKTHSVVKKHKQDRGEKKLLKCN 7F-1 7F-2 7F-3 DCEKIFSKISTLTLHQRIHTGEKPYECIECGKAFSQSAHLAQHORIHTGEKPFECTECGKAFSQNAHLVQHQRVHTG **ZF-4** ZF-5 EKPYQCKQCNKAFSQLAHLAQHQRVHTGEKPYECIECGKAFSDCSSLAHHRRIHTGKRPYECIDCGKAFRQNASLIR **ZF-7** ZF-8 7F-9 HRRYYHTGEKPFDCIDCGKAFTDHIGLIQHKRIHTGERPYKCNVCGKAFSHGSSLTVHQRIHTGEKPYECNICEKAF ZF-10 ZF-11 SHRGSLTLHQRVHTGEKPYECKECGKAFRQSTHLAHHQRIHTGEKPYECKECSKTFSQNAHLAQHQKIHTGEKPYEC 7F-12 ZF-13 ZF-14 KERGKAFSQIAHLVQHQRVHTGEKPYECIECGKAFSDGSYLVQHPRLHSGKRPYECLECGKAFRQRASLICHQRCHT ZF-15 ZF-16 ZF-17 GEKPYECNVCGKAFSHRKSLTLHQRIHTGEKPYECKECSKAFSQVAHLTLHKRIHTGERPYECKECGKAFRQSVHLA HHQRIHTGESSVILSSALPYHQVL* Fig. 4

KRAB-A MTDGLVTFRDVAIDFSQEEWECLDPAQRDLYVDVMLENYSNLVSLDLESKTYETKKIFSENDIFEINFSQWEMK DKSKTLGLEASIFRNNWKCKSIFEGLKGHQEGYFSQMIISYEKIPSYRKSKSLTPHQRIHNTEKSYVCKECGK ZF-2 ZF-3 ACSHGSKLVQHERTHTAEKHFECKECGKNYLSAYQLNVHQRFHTGEKPYECKECGKTFSWGSSLVKHERIHT **ZF-4** 7F-5 7F-6 GEKPYECKECGKAFSRGYHLTQHQKIHIGVKSYKCKECGKAFFWGSSLAKHEIIHTGEKPYKCKECGKAFSR ZF-7 **ZF-8** GYQLTQHQKIHTGKKPYECKICGKAFCWGYQLTRHQIFHTGEKPYECKECGKAFNCGSSLIQHERIHTGEKP **ZF-9** ZF-10 ZF-11 YECKECGKAFSRGYHLSQHQKIHTGEKPFECKECGKAFSWGSSLVKHERVHTGEKSHECKECGKTFCSGYQLT RHQVFHTGEKPYECKECGKAFNCGSSLVQHERIHTGEKPYECKECGRLLVVAITLLNIRKFIPYRNLSNVRNV GRPSVGVQA *

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Fig. 5A

Sequence of CZF-2 (cDNA)
50 GGGAGTTCTTGCAATTCCAGAACCATGACTGATGGGTTGGTGACATTCAG
100 GGATGTGGCCATCGACTTCTCAGGAGGAGTGGGAATGCCTGGACCCTG
150 CTCAGAGGGACTTGTACGTGGATGTAATGTTGGAGAACTATAGTAACTTG
200 GTGTCACTGGATTTGGAGTCAAAAACGTATGAGACCAAAAAATATTTTTC
250 AGAAAATGATATTTTTGAAATAAATTTTTCCCAGTGGGAGATGAAGGACA
300 AAAGTAAAACCCTTGGCCTTGAGGCATCCATCTTCAGAAATAATTGGAAG
350 TGCAAAAGCATATTCGAGGGACTAAAAGGACATCAAGAGGGATACTTCAG
400 TCAAATGATAATCAGCTATGAAAAAATACCTTCTTACAGAAAAAGTAAAT
450 CTCTTACTCCACATCAAAGAATTCATAATACAGAGAAATCCTATGTTTGT
500 AAGGAATGTGGGAAGGCTTGCAGTCATGGCTCAAAACTTGTTCAACATGA
550 GAGAACTCATACAGCTGAAAAGCACTTTGAATGTAAAGAATGTGGGAAGA
600 ATTATTTAAGTGCCTATCAACTCAATGTGCATCAGAGATTTCATACTGGT
650 GAGAAACCCTATGAGTGTAAGGAATGTGGGAAGACCTTTAGCTGGGGATC
700 AAGCCTTGTTAAACATGAGAGAATTCACACTGGTGAGAAACCCTATGAAT
750 GTAAAGAATGTGGGAAGGCCTTTAGTCGTGGCTATCACCTTACCCAACAT
800

 ${\tt CAGAAAATTCATATTGGTGTGAAATCTTATAAATGTAAGGAATGTGGGAA}$

Fig. 5B

850
GGCCTTTTTTTGGGGCTCAAGCCTTGCTAAACATGAGATAATTCATACAG
900 GTGAGAAACCTTATAAATGTAAAGAATGTGGGAAGGCCTTCAGTCGTGGC
GIGAGAAACCIIAIAAAIGIAAAGAAIGIGGGAAGGCCTTCAGTCGTGGC
950
TATCAACTTACTCAGCATCAGAAAATCCATACTGGTAAGAAACCTTATGA
1000
ATGTAAAATATGTGGAAAGGCTTTTTGTTGGGGCTATCAACTTACTCGAC
1050
ATCAGATATTCATACTGGTGAGAAACCCTATGAATGCAAGGAATGTGGG
1100
AAGGCTTTTAATTGCGGATCAAGTCTTATTCAACATGAAAGAATTCATAC
1150
TGGTGAGAAACCTTATGAATGTAAAGAATGTGGAAAGGCCTTTAGTCGTG
1200
GCTATCACCTTTCTCAACATCAGAAAATCCATACTGGTGAGAAACCTTTT
1250
GAATGTAAGGAATGTGGGAAGGCCTTTAGTTGGGGTTCAAGCCTTGTTAA
1300
ACATGAGAGAGTTCATACTGGTGAGAAATCCCATGAATGTAAAGAATGCG
1350
GAAAGACCTTTTGTAGTGGGTATCAACTTACTCGACATCAGGTATTTCAC
1400
ACTGGTGAGAAACCCTATGAATGTAAGGAATGTGGGAAGGCTTTTAATTG
1450
TGGATCAAGCCTTGTTCAACATGAAAGAATCCATACAGGGGAGAAACCCT
1500
ATGAATGTAAAGAATGTGGAAGGCTTTTAGTCGTGGCTATCACCTTACTC
1550
AACATCAGAAAATTCATACCGGTGAGAAACCTTTCAAATGTAAGGAATGT
1600
GGGAAGGCCTTCAGTTGGGGTTCAAGCCTAGTTAAGCATGAGAGAGTCCA

Fig. 5C

1650
TACTAATGAGAAGTCTTATGAATGTAAAGACTGTGGGAAGGCCTTTGGTA
1700 GTGGCTATCAACTTAGTGTTCATCAGAGATTTCATACTGGTGAGAAGCTT
1750 TATCAACATAAGGAATTCGGGAAGACCTTTACTCGTGGCTCAAAACTTGT
1800
TCATGAGAGAACTCATAGTAATGATAAACCCTACAAATATAACGAATGTG
1850
GGGAAGCCTTTCTGTGGACAACTTACTCAAATGAGAAAATTGATACTGAT
1900 GAAACCTTATGATTGAAAGTTGTAAAAGAATATTTTGTGTGTG
1950 ACAACTTATCATAATAAGAACTCTTACTCTTGAGAAACCTTGTGAATGTA
2000
AGGGTTGTGCAAAAGCCATTCATTTCTGTTTATGGGCAATTATCTTGCTA
2050
TCCAGCAATTCATACTAGTGAGAAATATTTTTGAATATAATTAAT
2100
AGGCCTTTAGACTTCTGTACAGTCTTATTGGATATCAATTTATACTGATG
2143 TAAAATCATTTAAATGAAAAAAAAAAAAAAAAAAAAAAA

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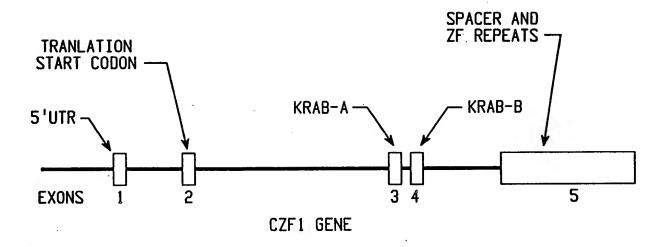
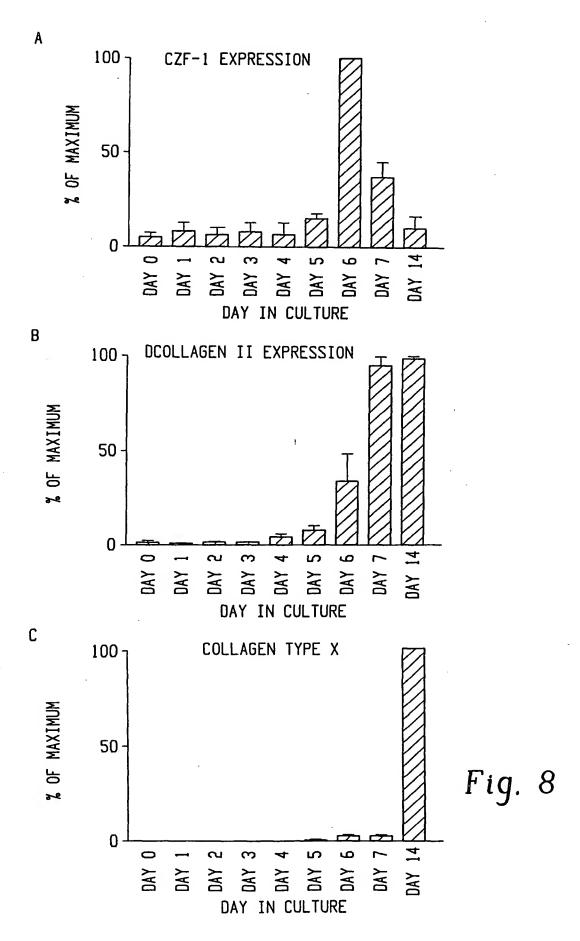
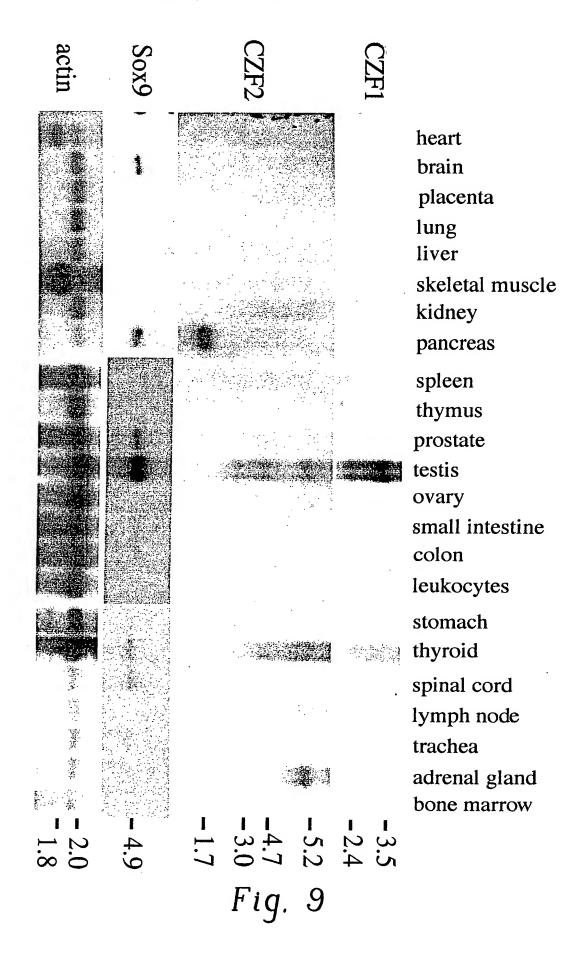


Fig. 7

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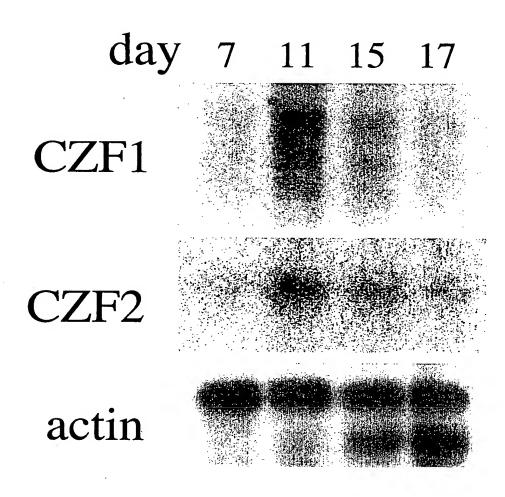
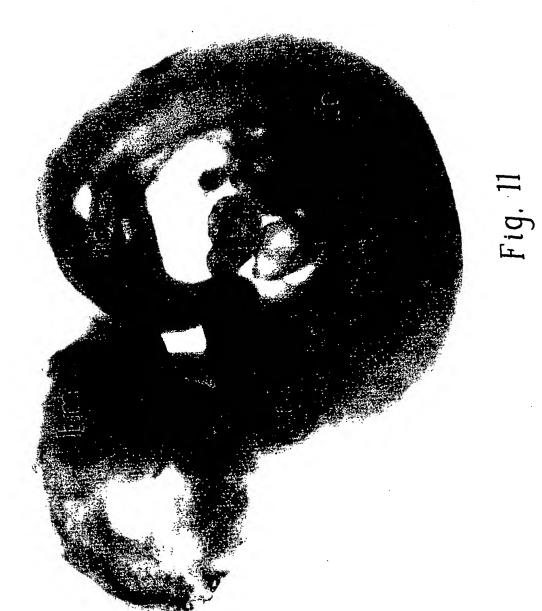


Fig. 10



forelimb bud (fb) tail bud (tb) head mesenchyme (hm) second branchial arch (b2) first branchial arch (b1)